

FIG. 1

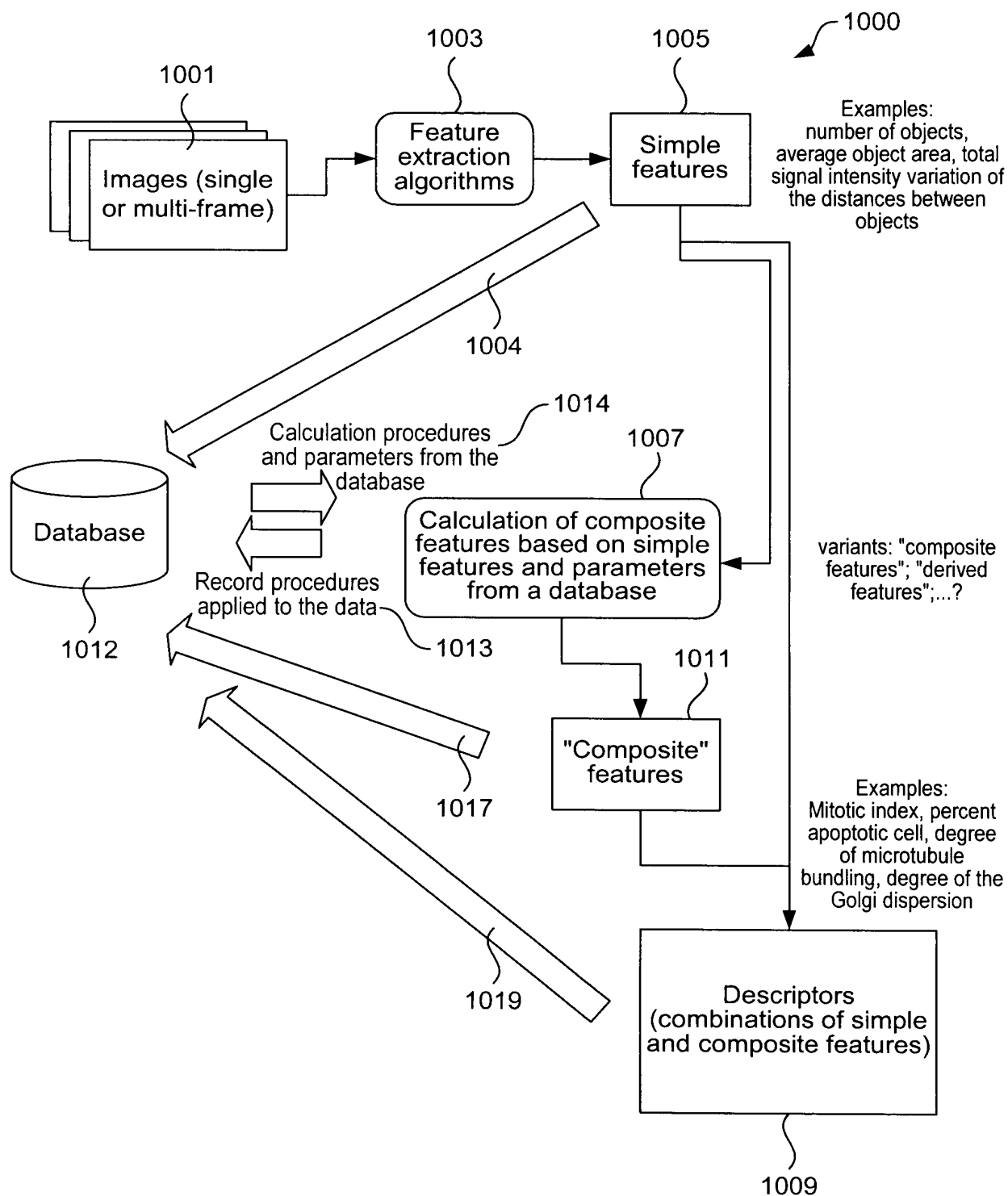


FIG. 1A

APPROVED	O. G. FIG.
BY	CLASS SUBCLASS
DRAFTSMAN	

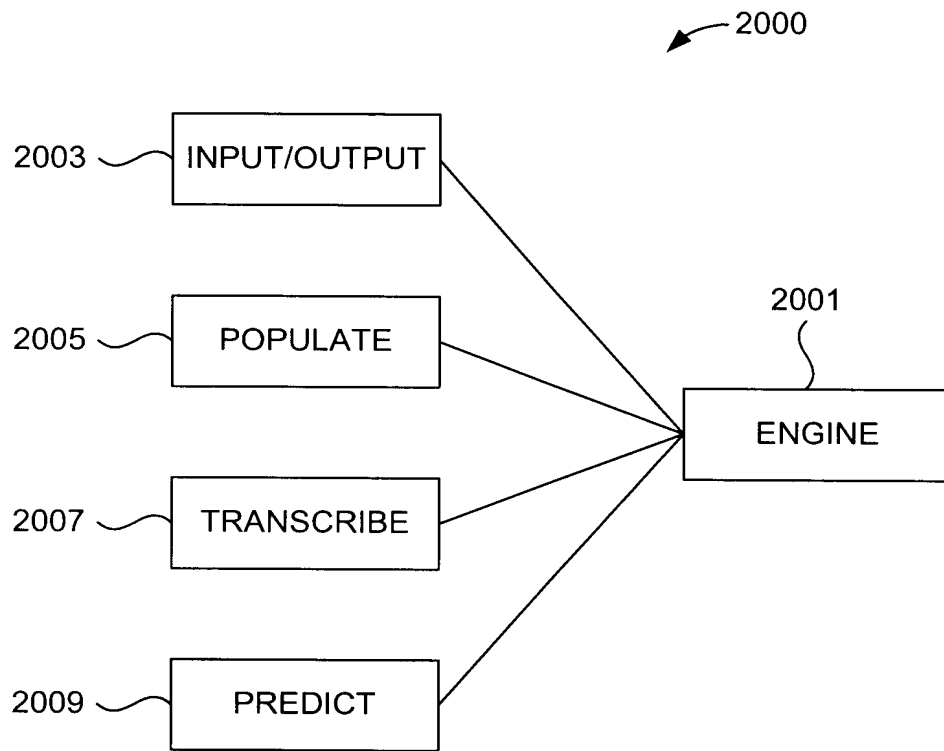


FIG. 1B

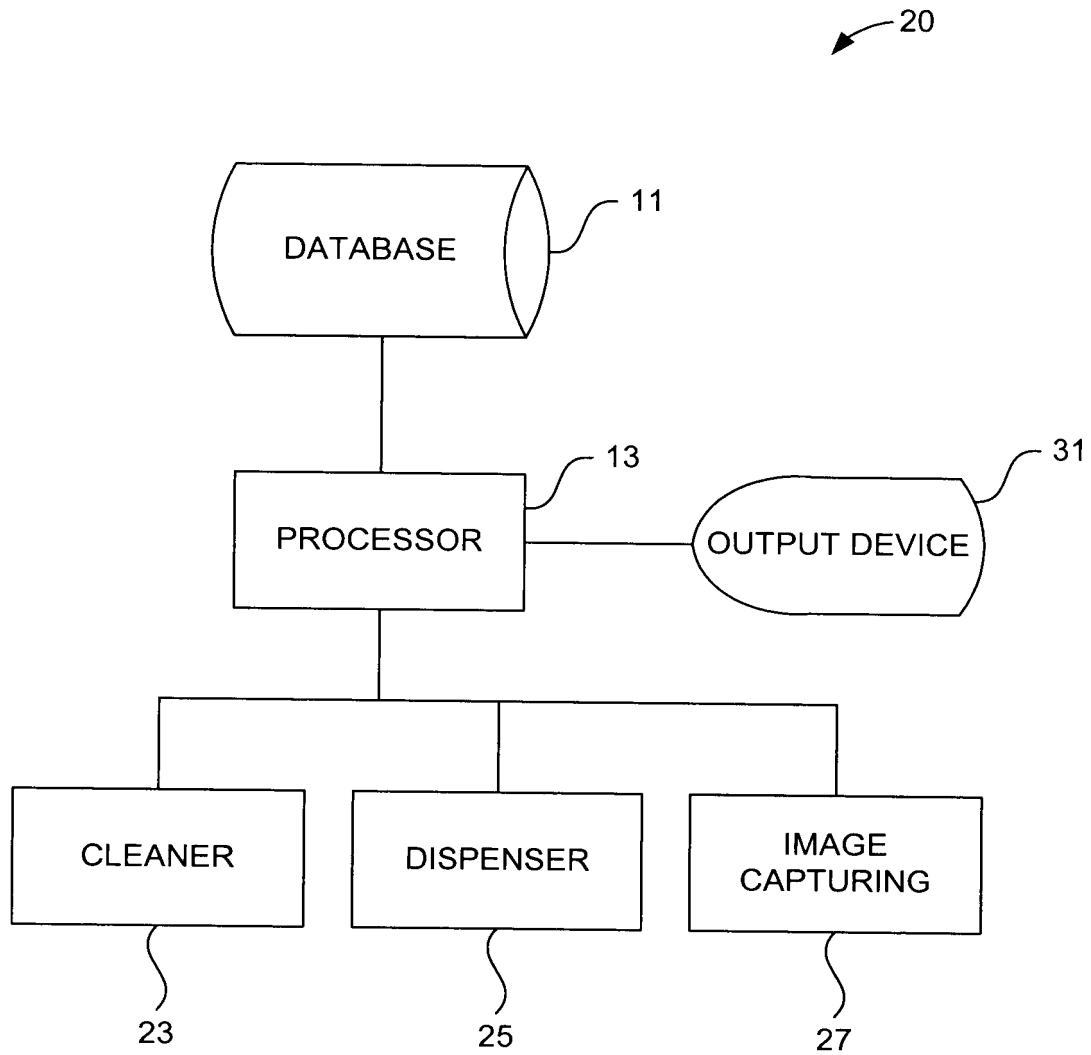


FIG. 2

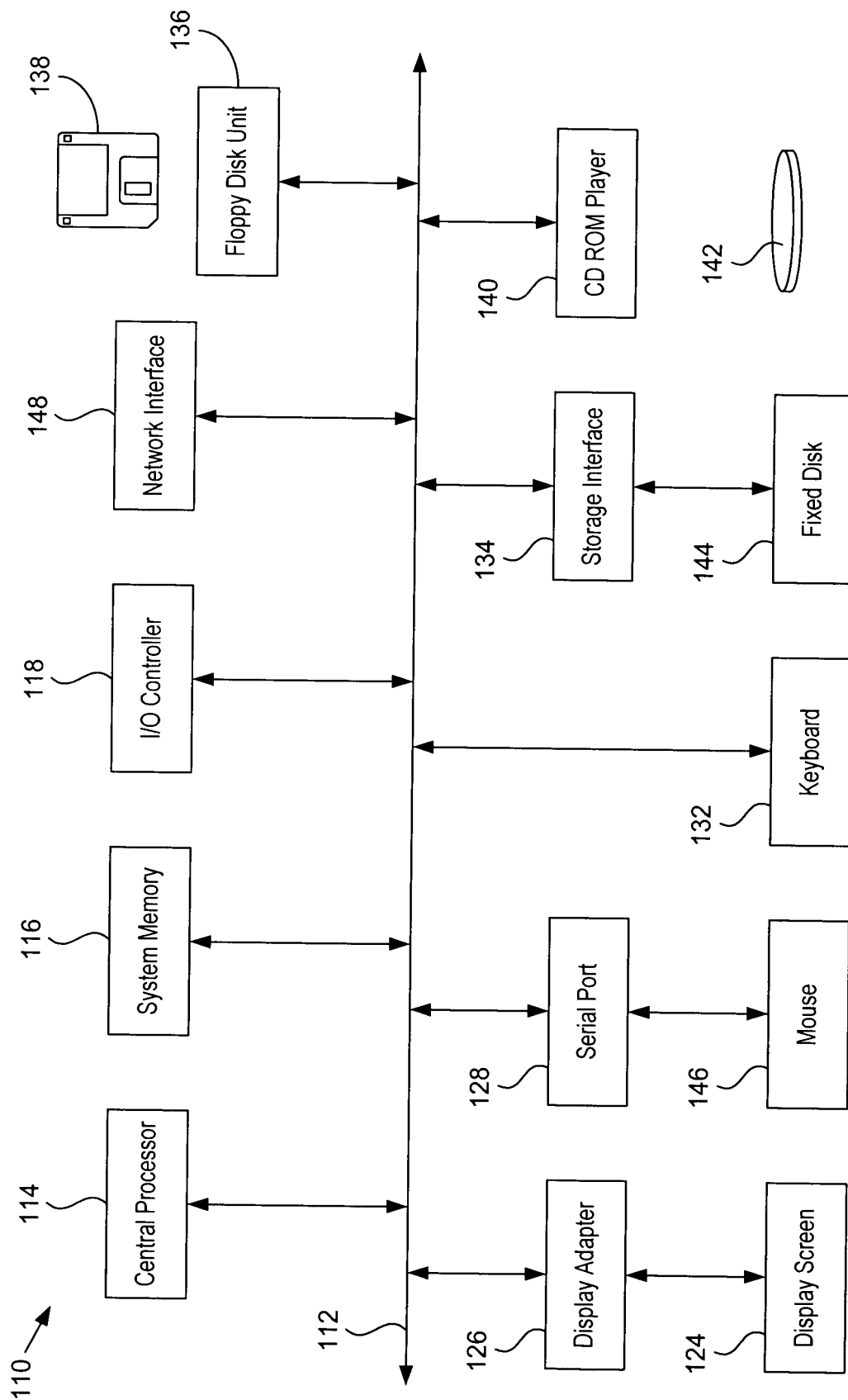


FIG. 3

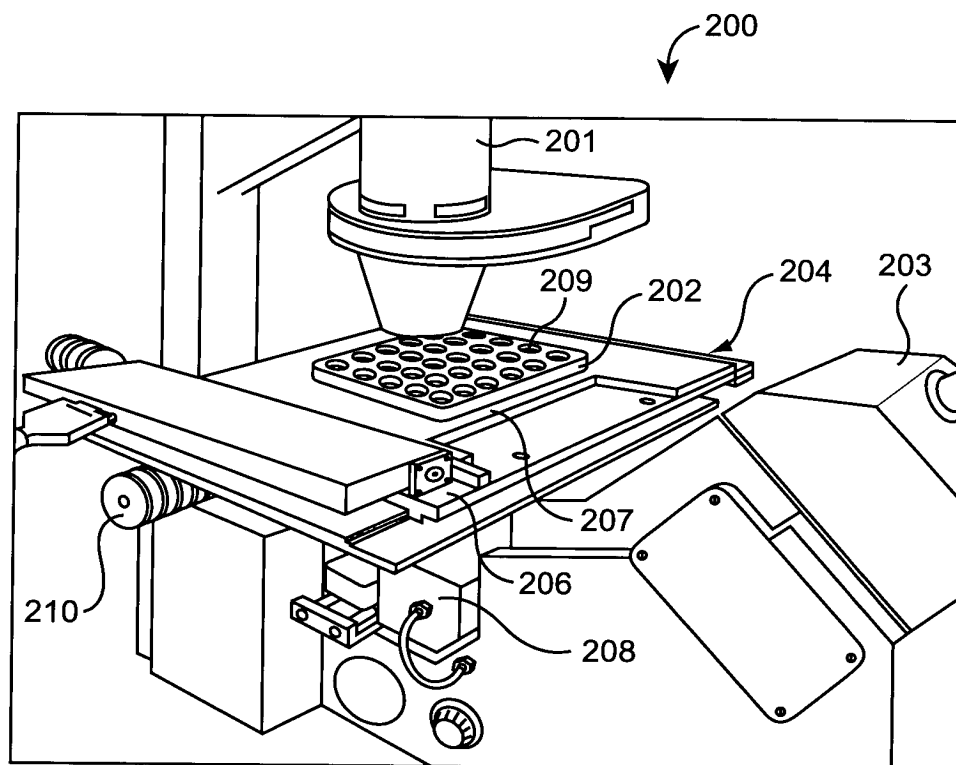


FIG. 4

APPROVED	Q. G. FIG.
BY	CLASS SUBCLASS
DRAFTSMAN	

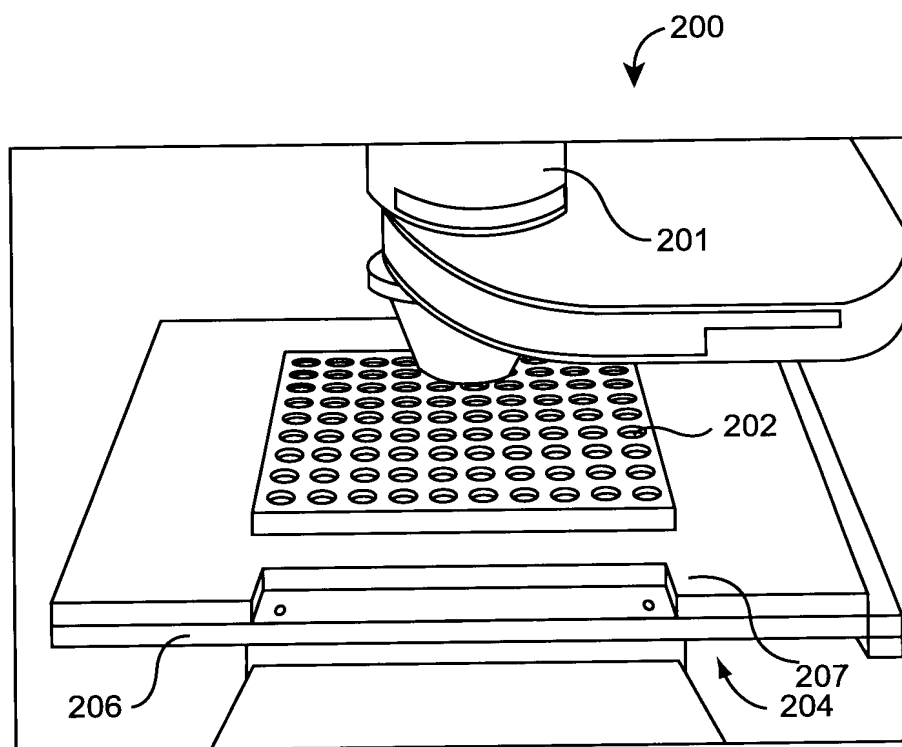


FIG. 5

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

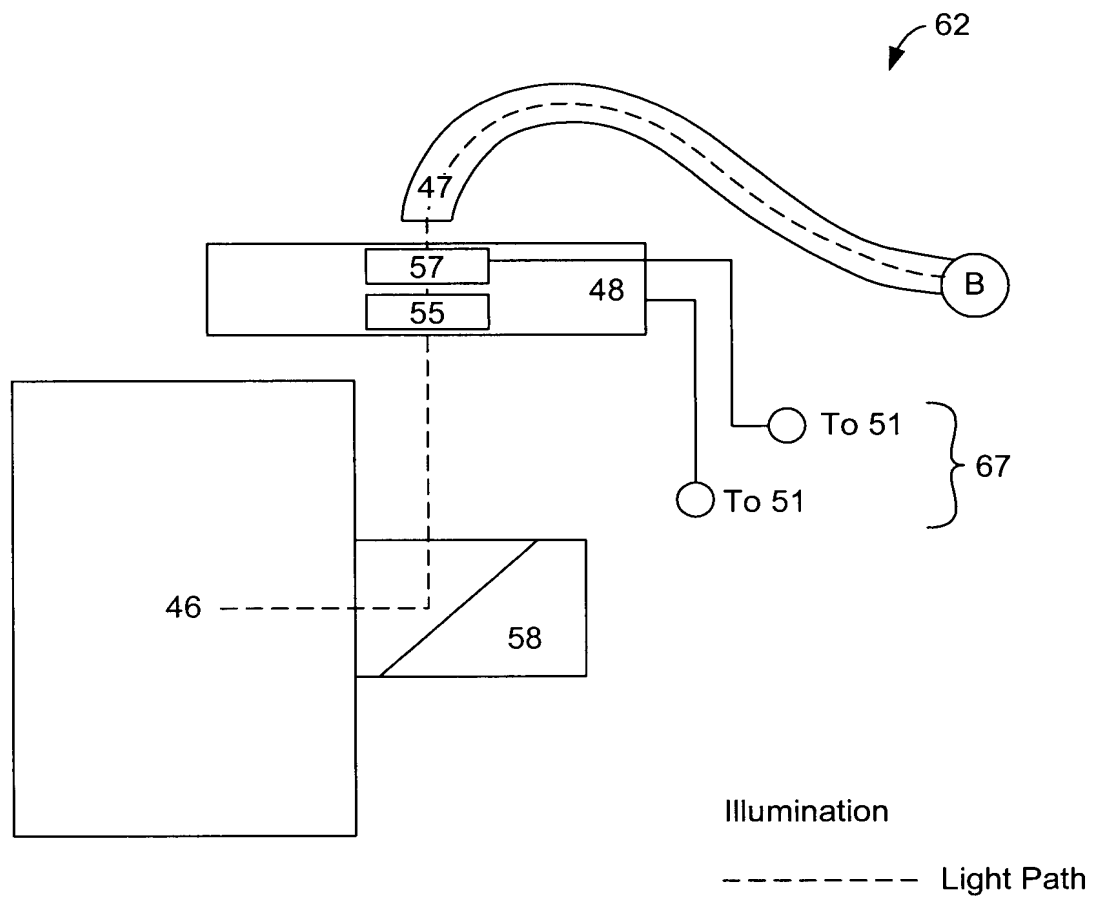


FIG. 5B

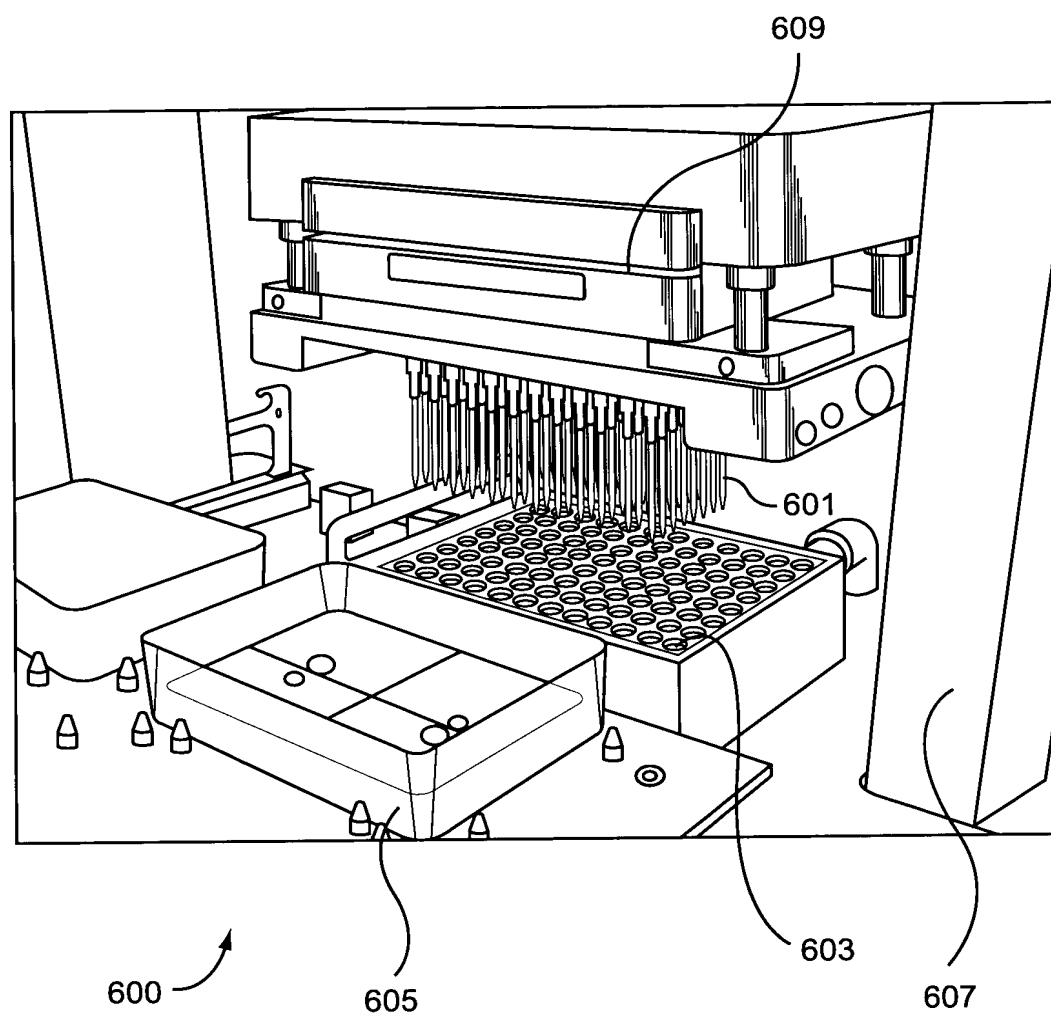


FIG. 6

APPROVED	O. G. FIG.
BY	CLASS/SUBCLASS
DRAFTSMAN	

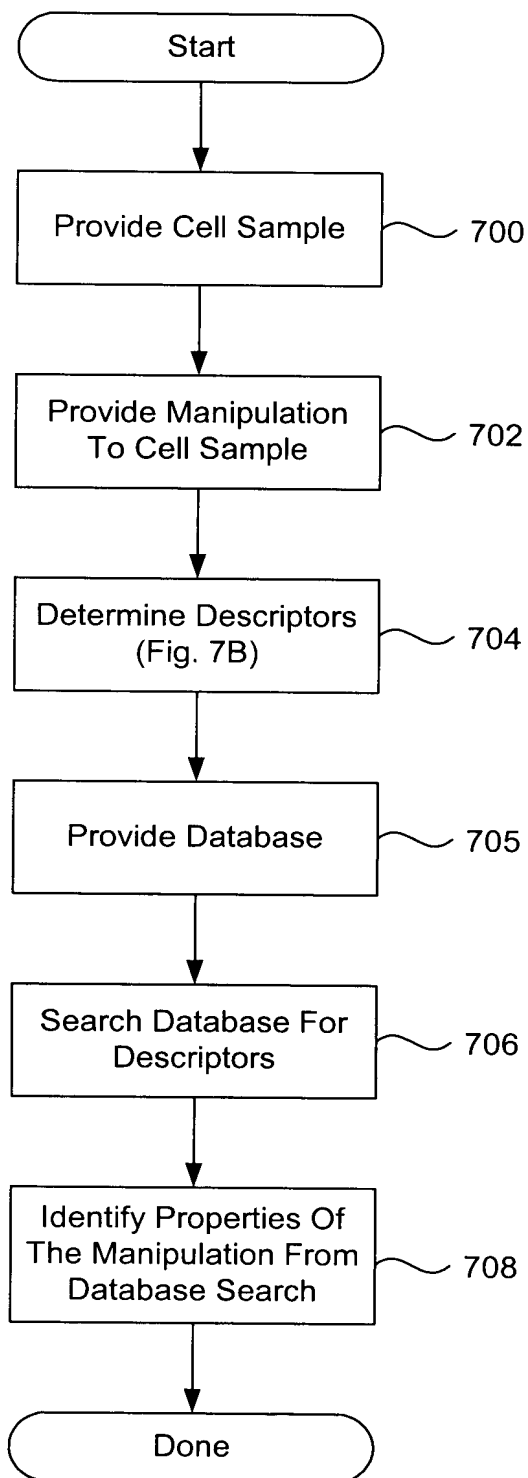


FIG. 7A

APPROVED	C. G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

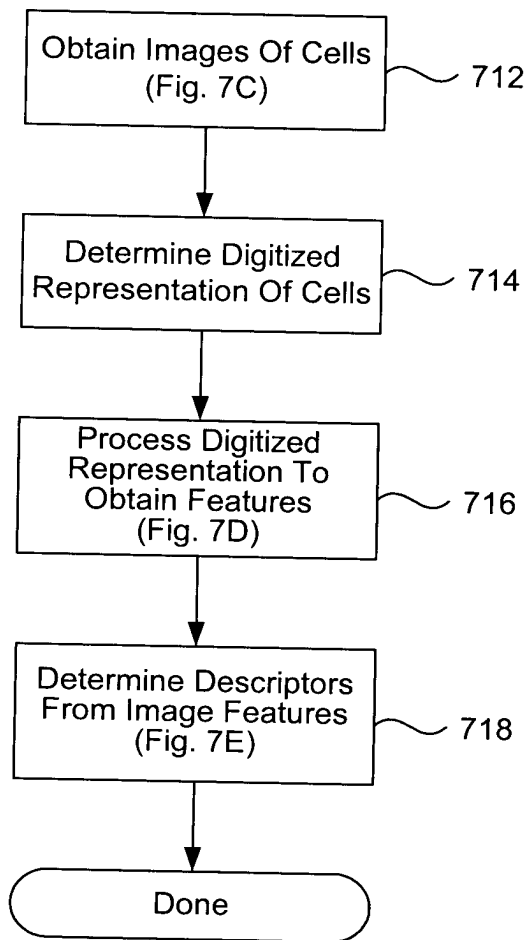


FIG. 7B
Step 704 of Fig. 7A

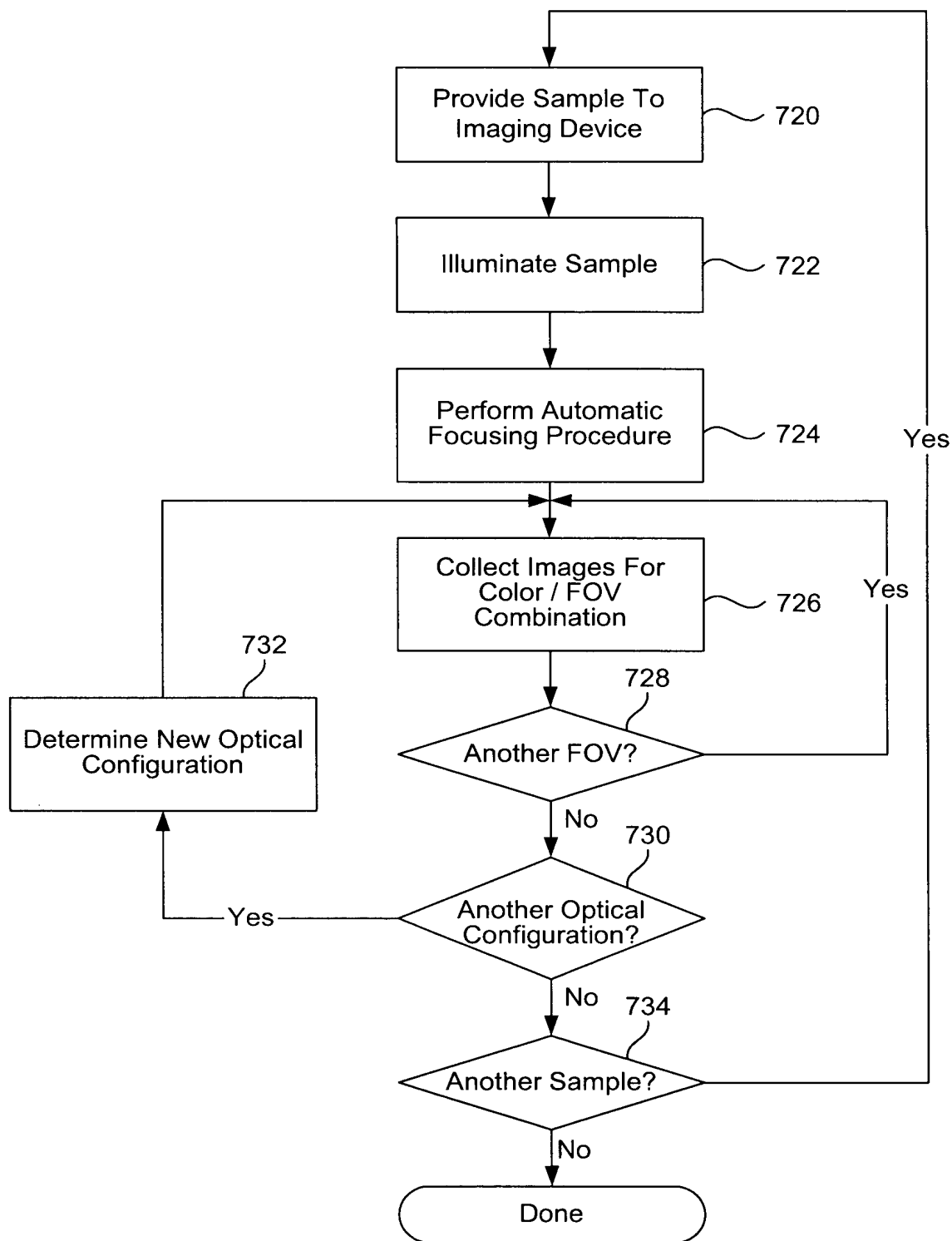


FIG. 7C
Step 714 of Fig. 7B

APPROVED	O G. FIG.
BY	CLASS SUBCLASS
DRAFTSMAN	

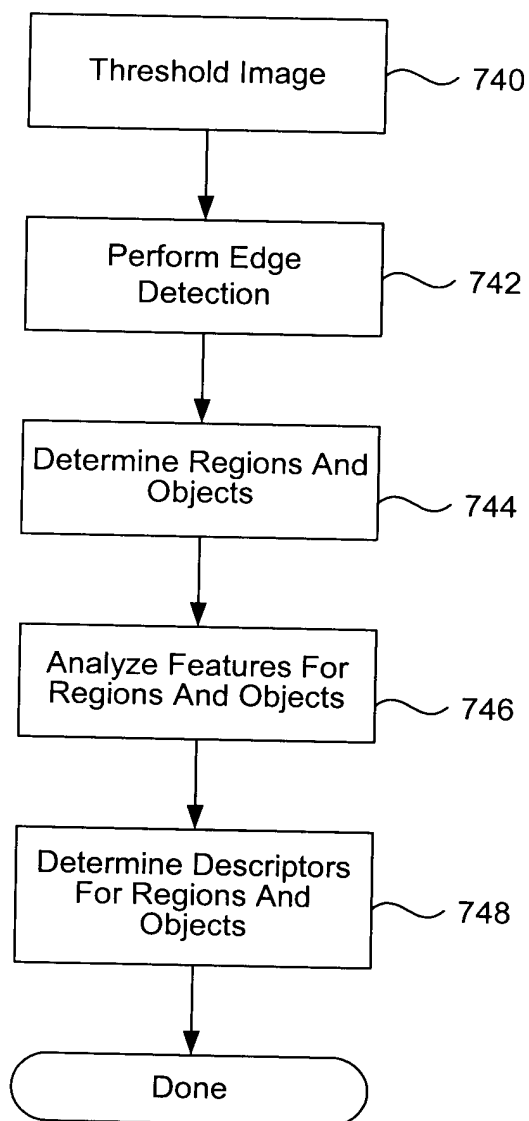


FIG. 7D
Step 716 of Fig. 7B

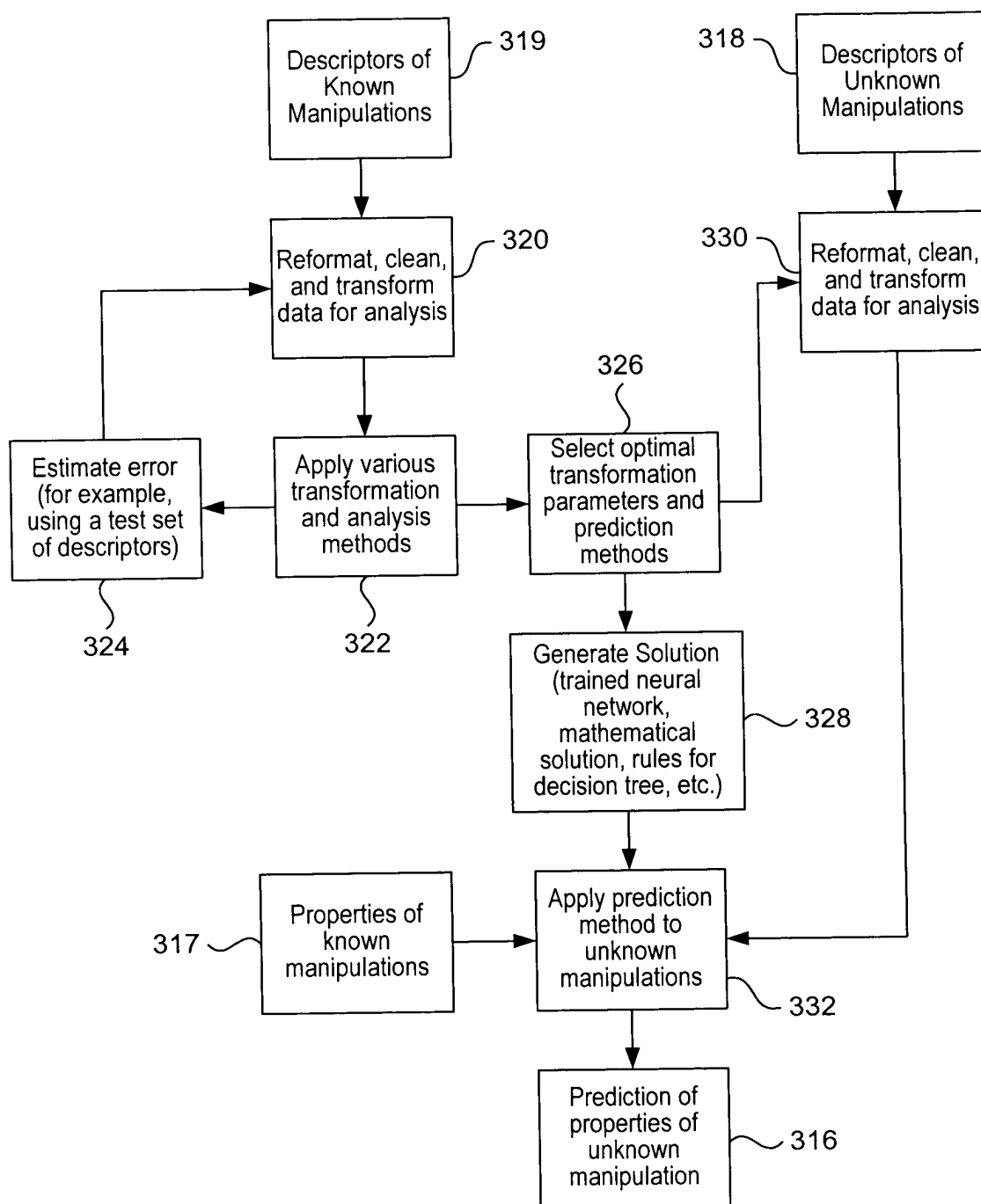


FIG. 7E

APPROVED	O. G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

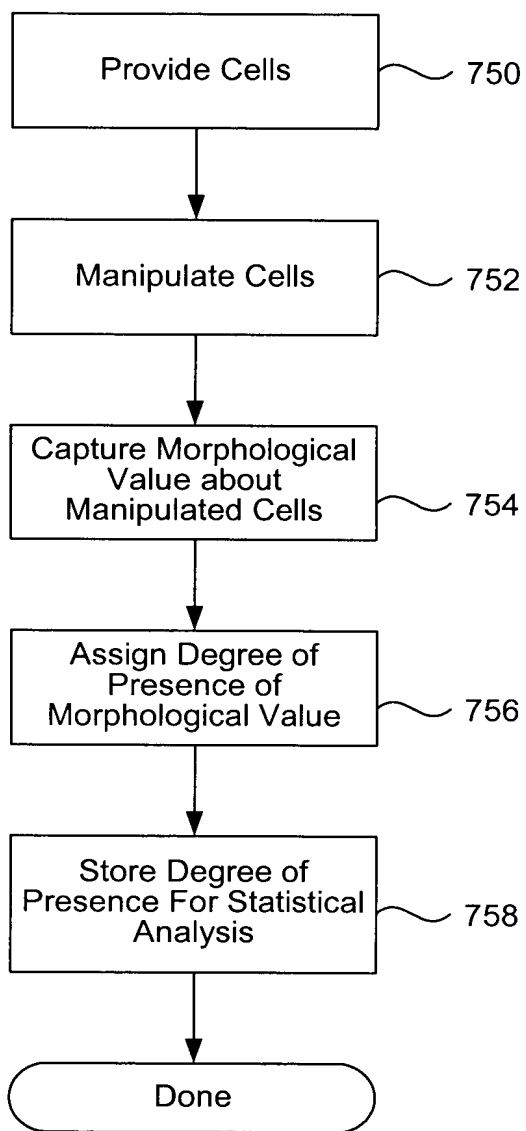


FIG. 7F

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

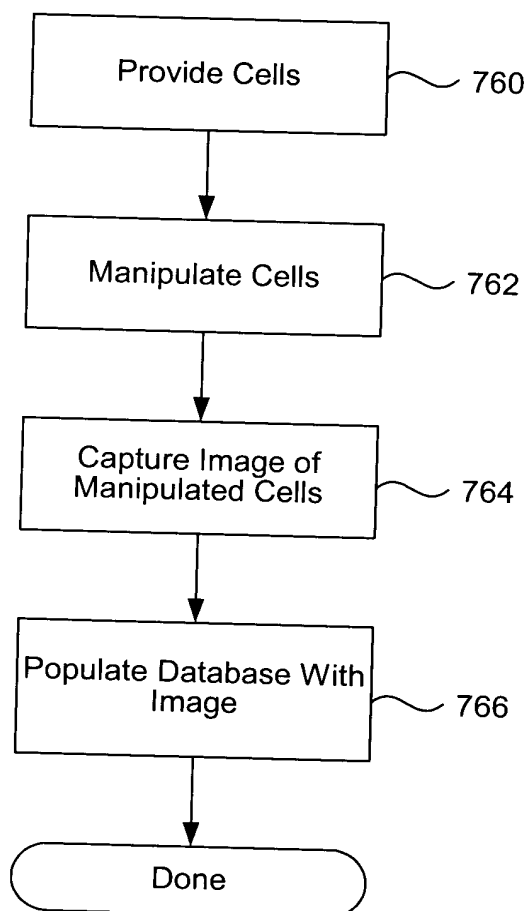


FIG. 7G

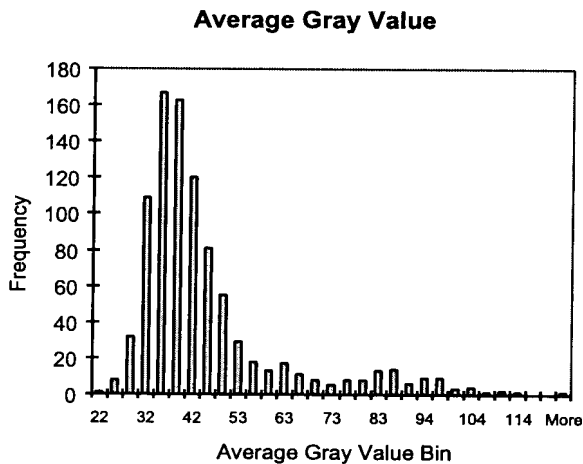


FIG. 8A

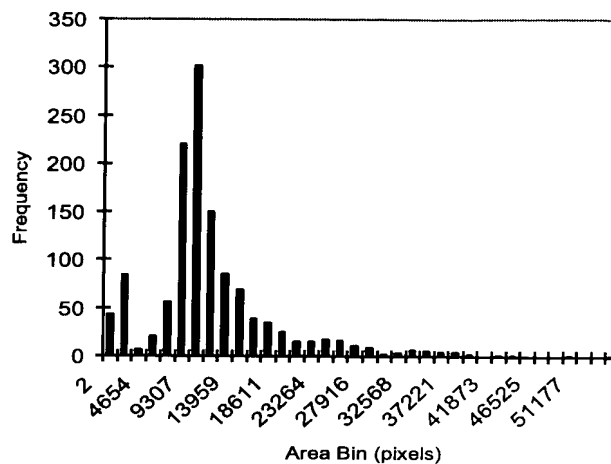


FIG. 8B

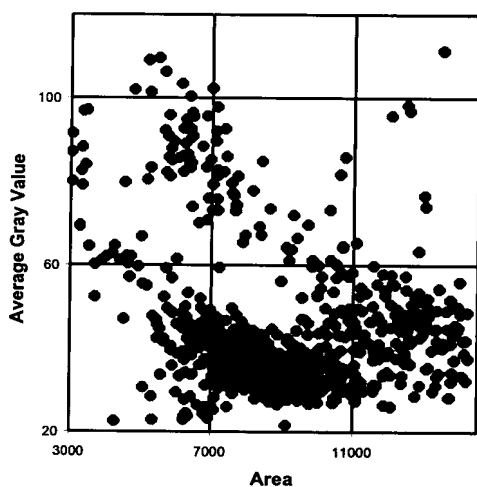


FIG. 8C

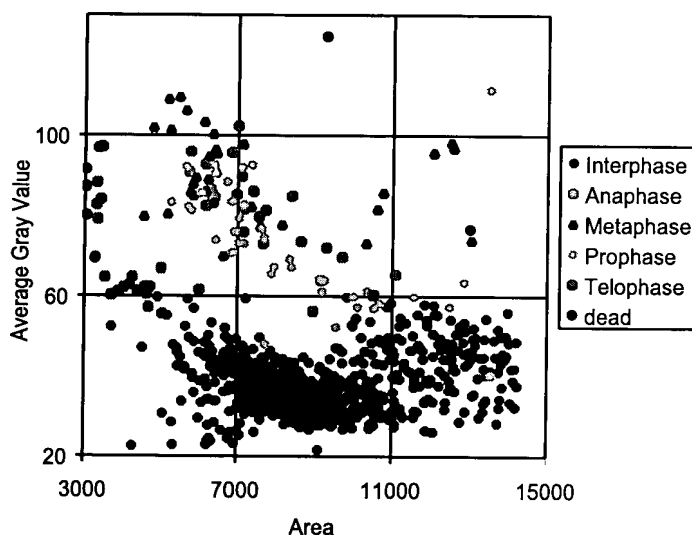


FIG. 8D

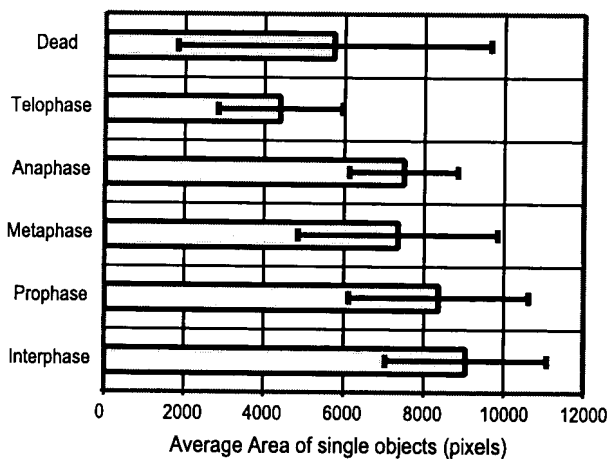


FIG. 8E

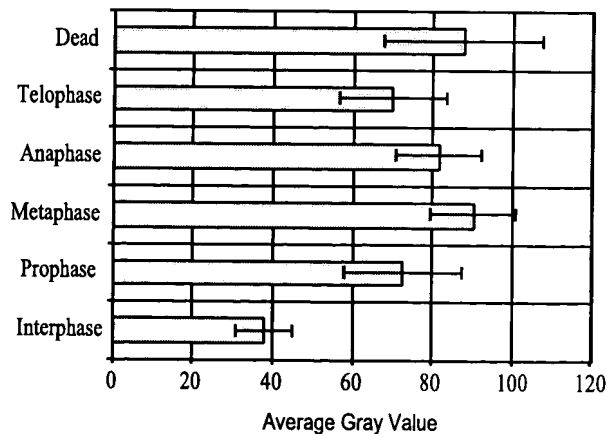


FIG. 8F

APPROVED	Q. G. FIG.
BY	CLASS SUBCLASS
DRAFTSMAN	

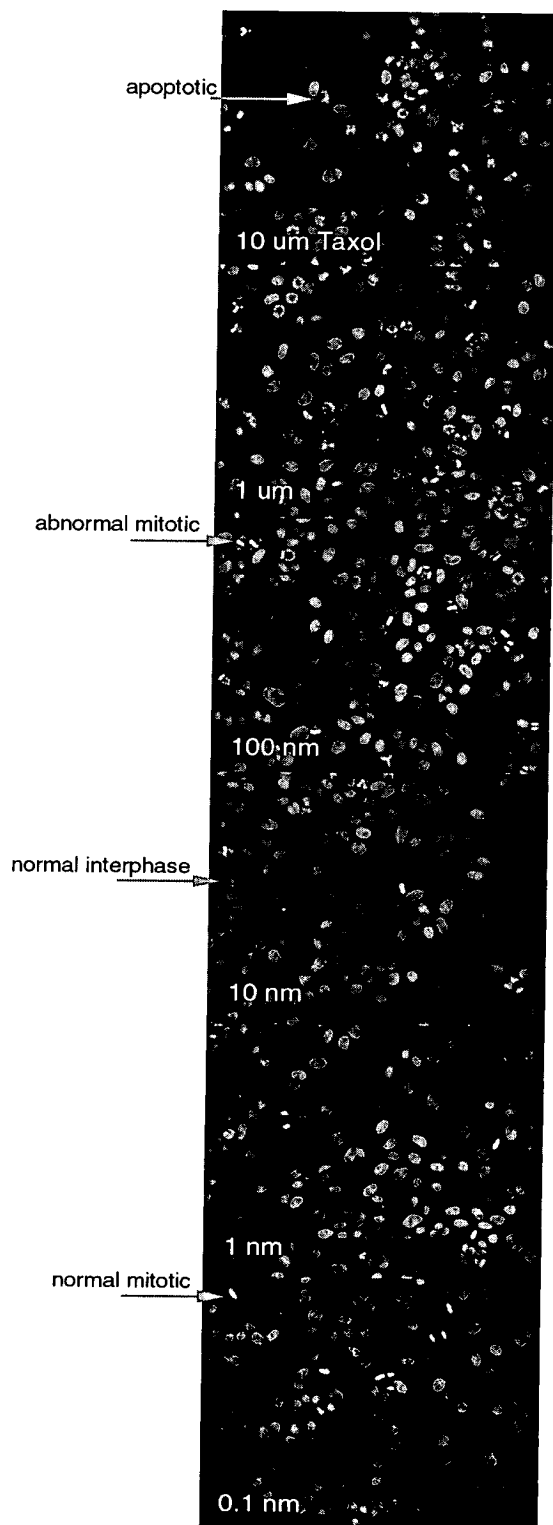


FIG. 9

MDCK cells treated with Taxol for 4.5 hours

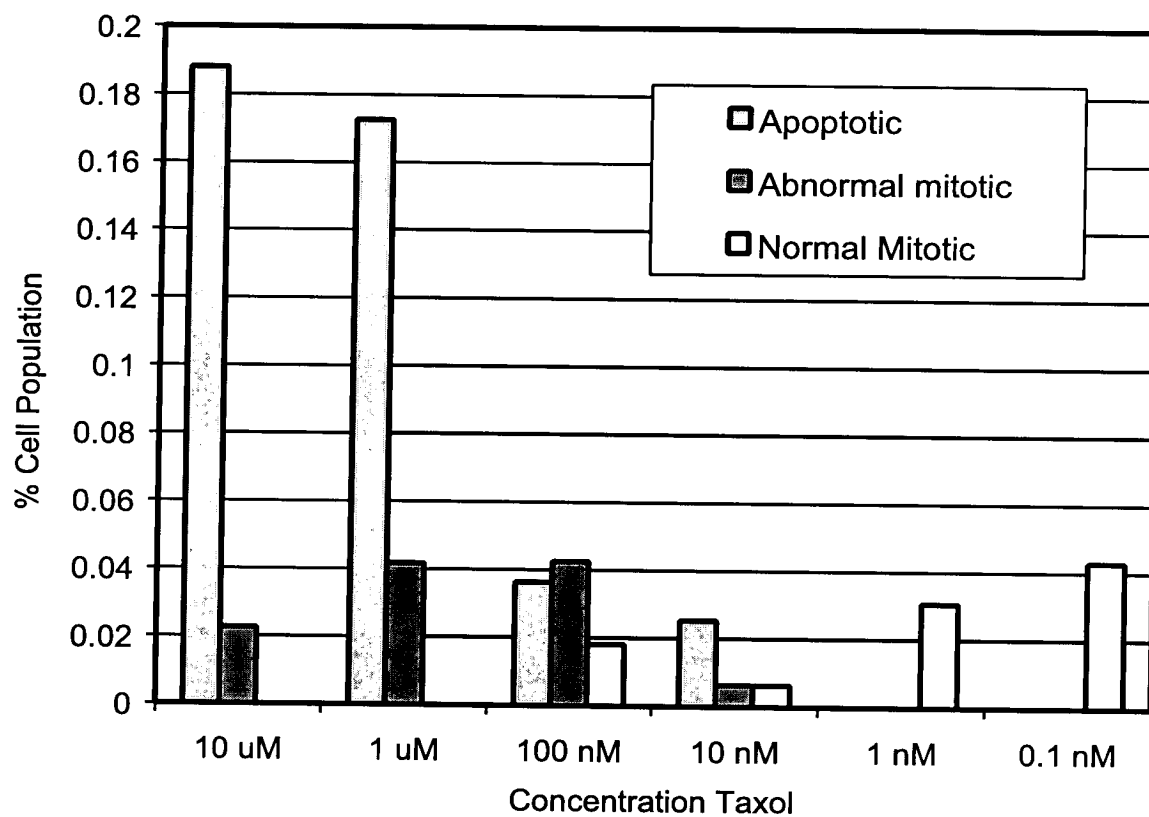


FIG. 10

APPROVED	O. G. FIG.
BY	CLASS/SUBCLASS
DRAFTSMAN	

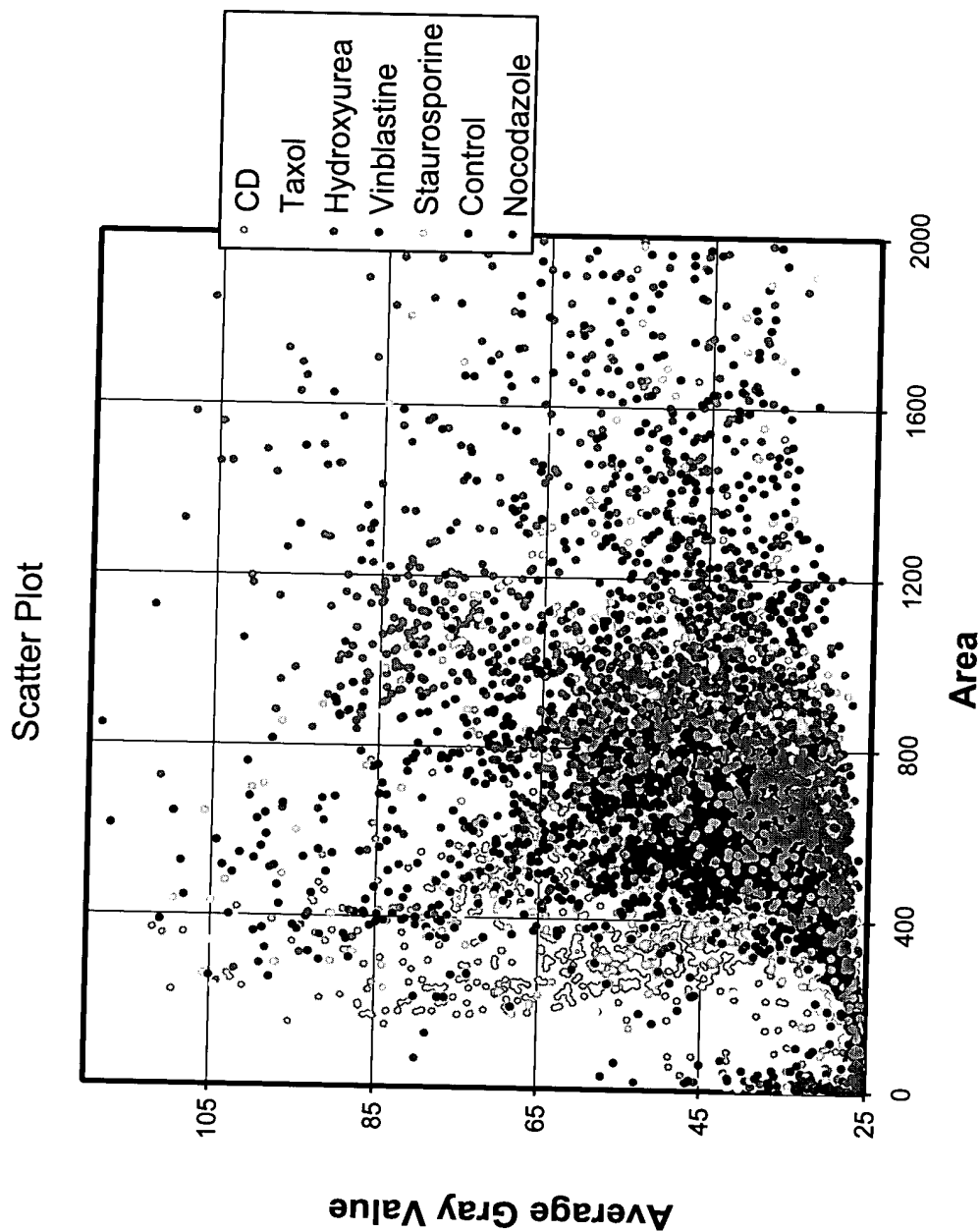


FIG. 11

APPROVED	O. G. FIG.	
BY	CLASS/SUBCLASS	
DRAFTSMAN		

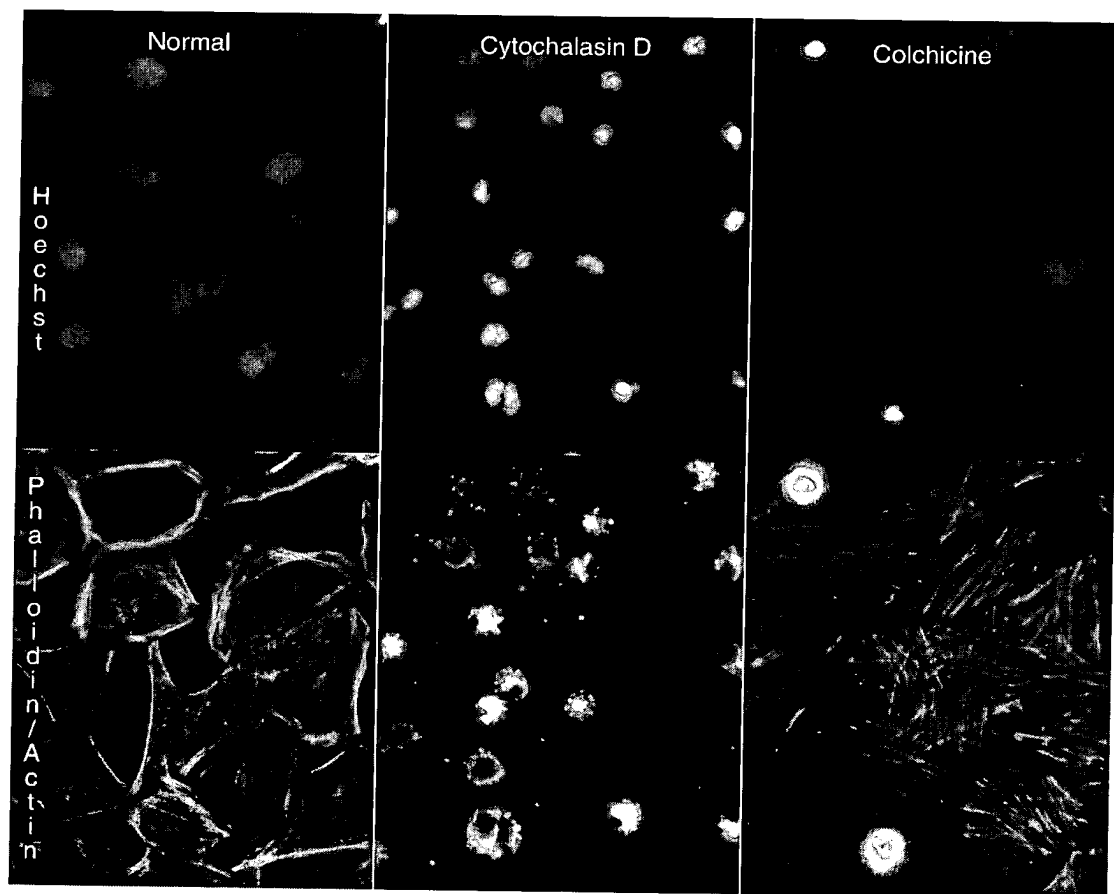


FIG. 12

APPROVED	O. G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

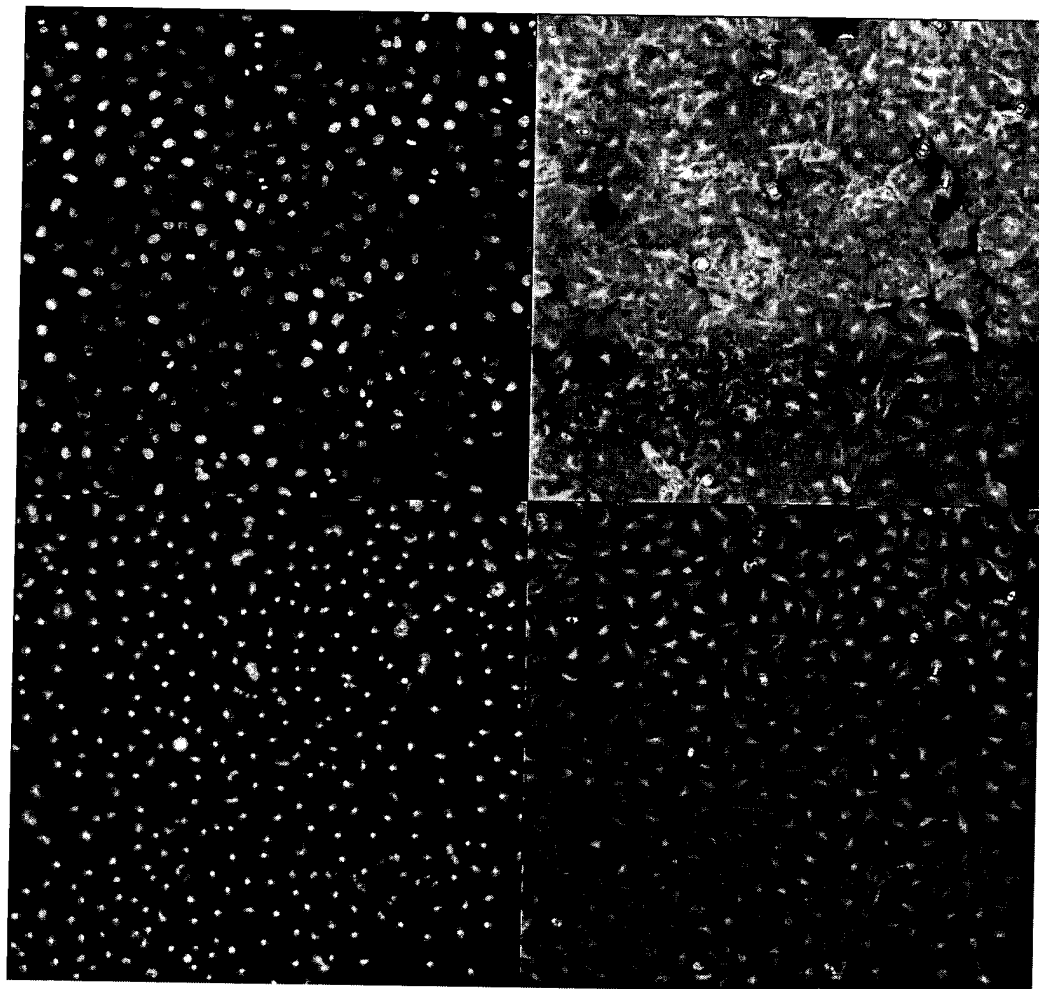


FIG. 13

APPROVED	O G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

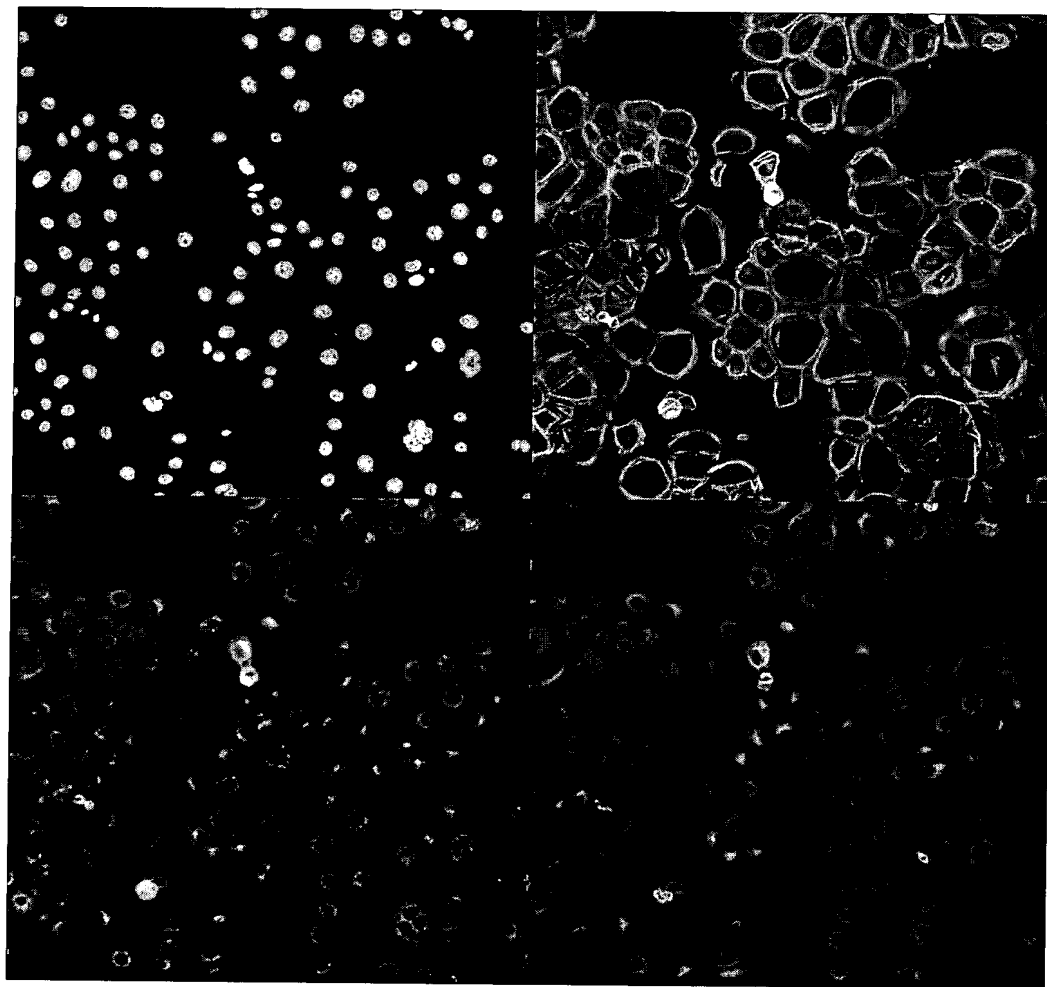


FIG. 14

APPROVED	O. G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

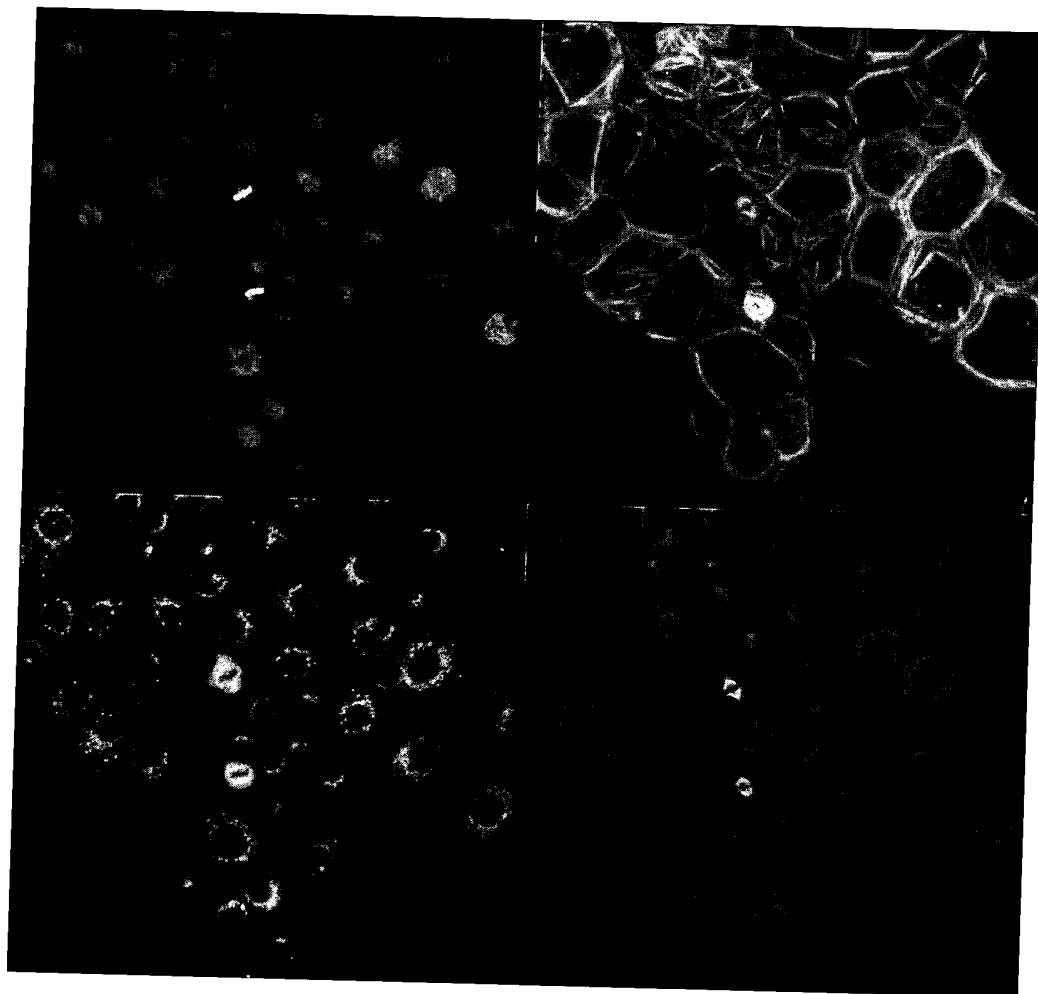


FIG. 15

APPROVED	O G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

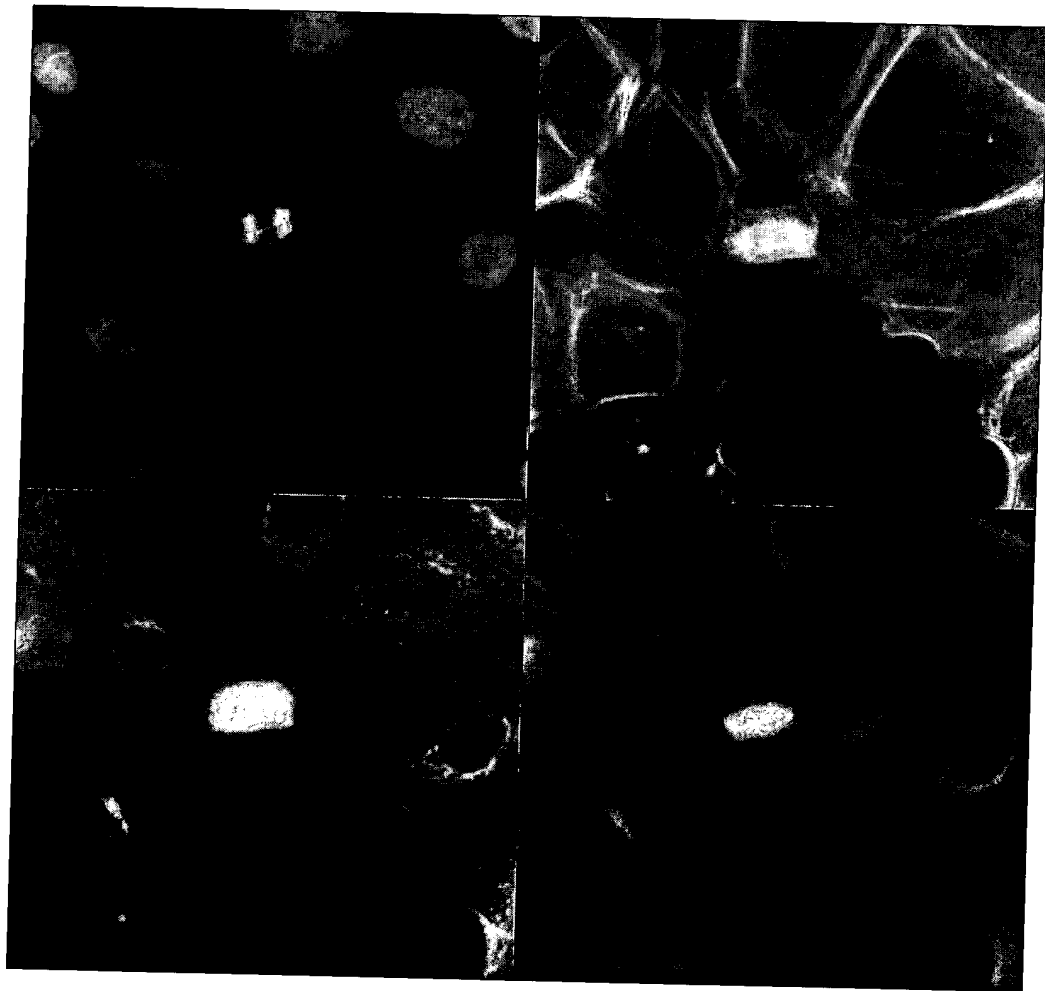


FIG. 16

APPROVED	O. G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

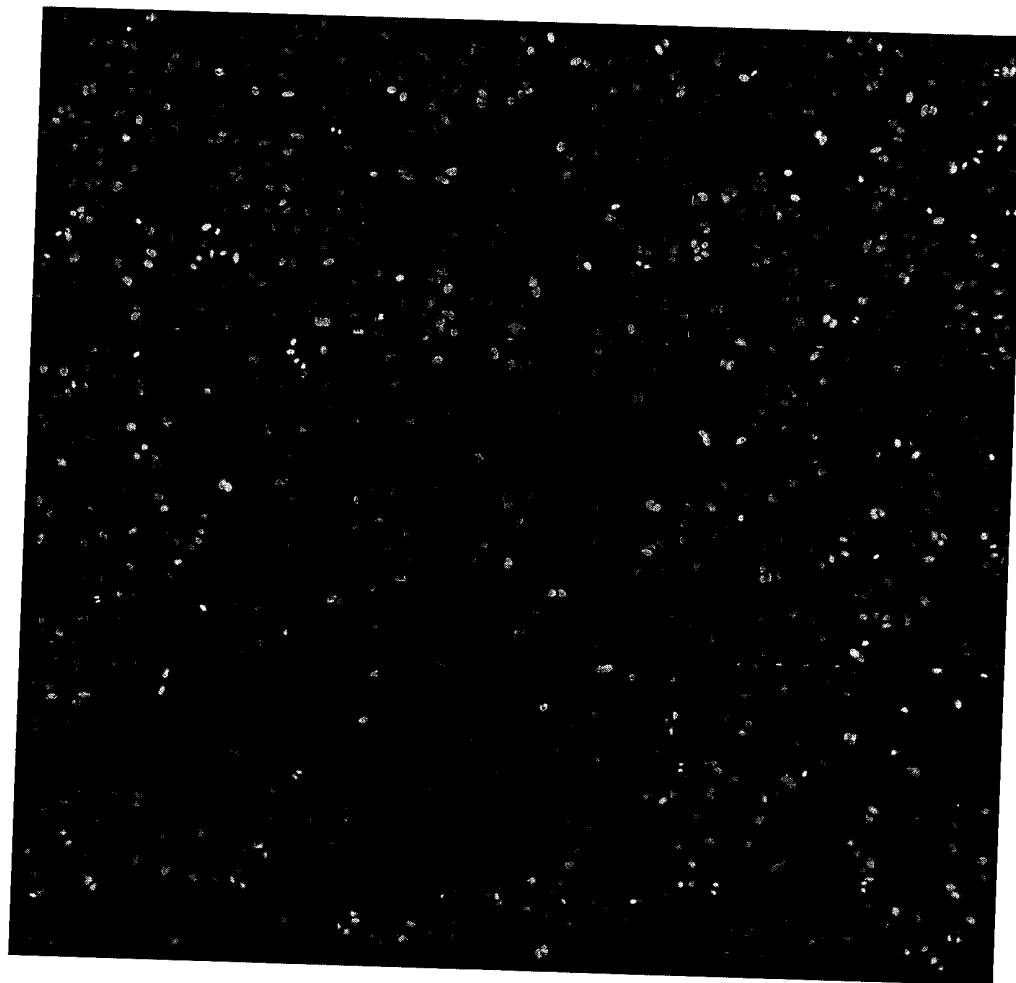


FIG. 17

APPROVED	O. G. FIG.
BY	CLASS SUBCLASS
DRAFTSMAN	

Conversion of morphometric parameters into nucleic acid code and clustering of the resulting sequences using Neighbor Joining method.

Compound:	Measurements																							
	Count	Area	Perimeter	Length	Breadth	Fiber length	Fiber breadth	Shape factor	Ell. form factor	Inner radius	Outer radius	Mean radius	Equiv. radius	Equiv. sphere vol.	Equiv. prolate vol.	Equiv. oblate vol.	Equiv. sphere surface area	Average gray value	Total gray value	Optical density	Radial dispersion	Texture Difference Moment	EFA Harmonic 2, Semi-Major	EFA Harmonic 2, Semi-Minor
Control	t	t	t	t	t	t	t	t	t	t	t	t	t	t	t	t	t	t	t	t	a	t	t	
Taxol	a	t	t	t	t	t	t	t	a	t	t	t	t	t	t	t	t	t	t	t	a	t	t	
CD	c	a	a	a	t	a	t	t	c	a	a	a	a	a	a	a	a	a	a	a	a	a	g	
Nocodozol	c	t	t	t	t	t	t	t	t	t	t	t	t	t	t	t	t	a	a	a	t	a	g	
Staurosporine	g	g	c	a	a	t	a	a	t	g	a	a	a	t	g	g	g	a	a	t	a	t	a	
Vinblastine	c	t	t	t	t	t	t	t	t	t	t	t	t	t	t	t	t	a	t	a	t	a	a	
Hydroxyurea	g	t	t	t	t	t	t	g	t	t	t	t	t	t	t	t	t	g	t	t	t	t	t	

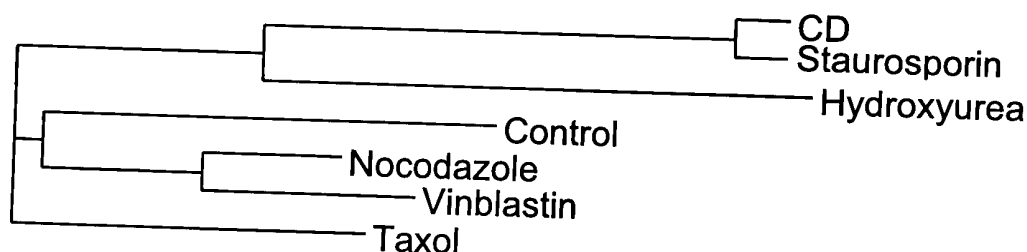


FIG. 18

APPROVED	O. G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

Conversion of morphometric parameters into amino acid codes and clustering of the resulting sequences using Neighbor Joining method.

	Count	Area	Perimeter	Length	Breadth	Fiber length	Fiber breadth	Shape factor	Ell. form factor	Inner radius	Outer radius	Mean radius	Equiv. radius	Equiv. sphere vol.	Equiv. prolate vol.	Equiv. oblate vol.	Equiv. sphere surface area	Average gray value	Total gray value	Optical density	Radial dispersion	Texture Difference Mome	EFA Harmonic 2, Semi-Major Axis	EFA Harmonic 2, Semi-Minor Ax	EFA Harmonic 2, Semi-Major A
Control	H	P	T	T	N	S	D	W	E	S	T	T	T	F	C	C	P	P	M	C	T	G	T	T	Y
Taxol	G	F	M	M	P	M	P	H	G	S	M	M	W	C	F	P	F	R	C	M	M	H	M	P	S
CD	F	G	G	G	M	G	M	K	A	G	G	G	G	G	G	G	G	H	G	G	G	M	G	V	H
Nocodozol	W	F	M	M	W	M	P	T	R	S	M	M	M	F	M	W	F	M	M	R	M	M	M	F	G
Staurosporine	N	V	A	G	G	M	G	G	Y	V	G	G	G	M	V	V	V	G	G	H	G	M	G	G	V
Vinblastine	F	W	W	M	W	W	C	W	D	S	M	W	W	M	M	M	W	M	V	E	M	M	M	F	P
Hydroxyurea	S	H	H	H	H	H	H	V	H	H	H	H	H	H	H	H	H	H	A	H	G	H	H	D	

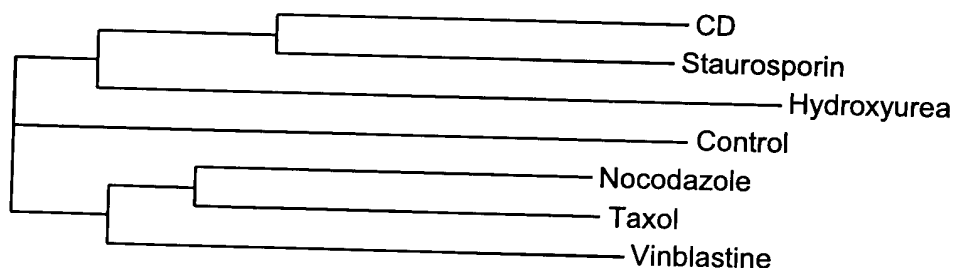


FIG. 19